

R. Schwabron

Re-run

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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/08/921,060B**

DATE: 01/25/2002  
TIME: 05:52:37

INPUT SET: S36682.raw

**This Raw Listing contains the General  
Information Section and up to the first 5 pages.**

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: ANDERSON, DARRELL R.  
HANNA, NABIL  
LEONARD, JOHN E.  
NEWMAN, ROLAND A.  
REFF, MITCHELL E.  
RASTETTER, WILLIAM H.

(ii) TITLE OF INVENTION: THERAPEUTIC APPLICATION OF CHIMERIC AND  
RADIOLABELED ANTIBODIES TO HUMAN B LYMPHOCYTE RESTRICTED  
DIFFERENTIATION ANTIGEN FOR THE TREATMENT OF B CELL  
LYMPHOMA

(iii) NUMBER OF SEQUENCES: 11

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: PILLSBURY WINTHROP  
(B) STREET: 1100 New York Avenue, N.W., Ninth FL.  
(C) CITY: Washington  
(D) STATE: DC  
(E) COUNTRY: USA  
(F) ZIP: 20005

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/921,060  
(B) FILING DATE: 29-AUG-1997  
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/149,099  
(B) FILING DATE: 03-NOV-1993

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US 07/978,891  
(B) FILING DATE: 13-NOV-1992

(viii) ATTORNEY/AGENT INFORMATION:

**ENTERED**

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/921,060BDATE: 01/25/2002  
TIME: 05:52:37

INPUT SET: S36682.raw

47 (A) NAME: Teskin, Robin L.  
48 (B) REGISTRATION NUMBER: 35,030  
49 (C) REFERENCE/DOCKET NUMBER: 037003-0275463  
50

51 (ix) TELECOMMUNICATION INFORMATION:  
52 (A) TELEPHONE: 202-861-3000  
53 (B) TELEFAX: 202-822-0944  
54  
55

56 (2) INFORMATION FOR SEQ ID NO:1:  
57

58 (i) SEQUENCE CHARACTERISTICS:  
59 (A) LENGTH: 27 base pairs  
60 (B) TYPE: nucleic acid  
61 (C) STRANDEDNESS: single  
62 (D) TOPOLOGY: linear  
63

64 (ii) MOLECULE TYPE: DNA (genomic)  
65  
66  
67

68 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
69

70  
71 GGGAGCTTGG ATCGATCCTC TATGGTT  
72

27

73 (2) INFORMATION FOR SEQ ID NO:2:  
74

75 (i) SEQUENCE CHARACTERISTICS:  
76 (A) LENGTH: 8540 base pairs  
77 (B) TYPE: nucleic acid  
78 (C) STRANDEDNESS: single  
79 (D) TOPOLOGY: linear  
80

81 (ii) MOLECULE TYPE: DNA (genomic)  
82  
83  
84

85 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:  
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89 AGGCCGAGGC GGCCTCGGCC TCTGCATAAA TAAAAAAAAT TAGTCAGCCA TGCATGGGGC 120  
90 GGAGAATGGG CGGAAC TGGG CGGAGTTAGG GGCGGGATGG GCGGAGTTAG GGGCGGGACT 180  
91 ATGGTTGCTG ACTAATTGAG ATGCATGCTT TGCATACTTC TGCTTGCTGG GGAGCCTGGG 240  
92 GACTTTCCAC ACCTGGTTGC TGAATAATTG AGATGCATGC TTTGCATACT TCTGCCTGCT 300  
93 GGGGAGCCTG GGGACTTTCC ACACCCCTAAC TGACACACAT TCCACAGAAT TAATTCCCCT 360  
94 AGTTATTAAT AGTAATCAAT TACGGGGTCA TTAGTTCATA GCCCATATAT GGAGTTCCGC 420  
95 GTTACATAAC TTACGGTAAA TGGCCCGCCT GGCTGACCGC CCAACGACCC CCGCCCATTTG 480  
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99 ATGACCTTAT GGGACTTTCC TACTTGGCAG TACATCTACG TATTAGTCAT CGCTATTACC 720

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PATENT APPLICATION US/08/921,060BDATE: 01/25/2002  
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126	GGCGCCCTGA	CCAGCGGCGT	GCACACCTTC	CCGGCTGTCC	TACAGTCCTC	AGGACTCTAC	2340
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140	CTCTCCCTGT	CTCCGGGTAA	ATGAGGATCC	GTAAACGGTT	ACCAACTACC	TAGACTGGAT	3180
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151	GCAGGAGCCA	GGGCAGAGCA	TATAAGGTGA	GGTAGGATCA	GTTGCTCCTC	ACATTTGCTT	3840
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171	TCTCTGCACA	GATAAGGACA	AACATTATTC	AGAGGGAGTA	CCCAGAGCTG	AGACTCCTAA	5040
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173	CGTAGAGCCA	CACCTTGATA	AGGGCCAATC	TGCTCACACA	GGATAGAGAG	GGCAGGAGCC	5160
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201	TGCTGGCGTT	TTTCCATAGG	CTCCGCCCCC	CTGACGAGCA	TCACAAAAT	CGACGCTCAA	6840
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203	CCCTCGTGCG	CTCTCCTGTT	CCGACCCTGC	CGCTTACCGG	ATACCTGTCC	GCCTTTCTCC	6960
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RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/921,060BDATE: 01/25/2002  
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208 AGTGGTGGCC TAACTACGGC TACACTAGAA GGACAGTATT TGGTATCTGC GCTCTGCTGA 7260  
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210 GTAGCGGTGG TTTTTTTGTT TGCAAGCAGC AGATTACGCG CAGAAAAAAA GGATCTCAAG 7380  
211 AAGATCCTTT GATCTTTTCT ACGGGGTCTG ACGCTCAGTG GAACGAAAAC TCACGTTAAG 7440  
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217 GAAGGGCCGA GCGCAGAAGT GGTCTGCAA CTTTATCCGC CTCCATCCAG TCTATTAATT 7800  
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220 CCCAACGATC AAGGCGAGTT ACATGATCCC CCATGTTGTG CAAAAAAGCG GTTAGCTCCT 7980  
221 TCGGTCCTCC GATCGTTGTC AGAAGTAAGT TGGCCGCGAGT GTTATCACTC ATGGTTATGG 8040  
222 CAGCACTGCA TAATTCTCTT ACTGTCATGC CATCCGTAAG ATGCTTTTCT GTGACTGGTG 8100  
223 AGTACTCAAC CAAGTCATTG TGAGAATAGT GTATGCGGCG ACCGAGTTGC TCTTGCCCGG 8160  
224 CGTCAATACG GGATAATACC GCGCCACATA GCAGAACTTT AAAAGTGCTC ATCATTGGAA 8220  
225 AACGTTCTTC GGGGCGAAAA CTCTCAAGGA TCTTACCGCT GTTGAGATCC AGTTCGATGT 8280  
226 AACCCACTCG TGCACCCAAC TGATCTTCAG CATCTTTTAC TTTACCAGC GTTCTGGGT 8340  
227 GAGCAAAAAC AGGAAGGCAA AATGCCGCAA AAAAGGGAAT AAGGGCGACA CGGAAATGTT 8400  
228 GAATACTCAT ACTCTTCCTT TTTCAATATT ATTGAAGCAT TTATCAGGGT TATTGTCTCA 8460  
229 TGAGCGGATA CATATTTGAA TGTATTTAGA AAAATAAACA AATAGGGGTT CCGCGCACAT 8520  
230 TTCCCCGAAA AGTGCCACCT 8540

## (2) INFORMATION FOR SEQ ID NO:3:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 9209 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: DNA (genomic)

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

246  
247 GACGTCGCGG CCGCTCTAGG CCTCCAAAAA AGCCTCCTCA CTACTTCTGG AATAGCTCAG 60  
248 AGGCCGAGGC GGCCTCGGCC TCTGCATAAA TAAAAAAAT TAGTCAGCCA TGCATGGGGC 120  
249 GGAGAATGGG CGGAAGTGGG CGGAGTTAGG GGCGGGATGG GCGGAGTTAG GGGCGGGACT 180  
250 ATGGTTGCTG ACTAATTGAG ATGCATGCTT TGCATACTTC TGCCTGCTGG GGAGCCTGGG 240  
251 GACTTTCCAC ACCTGGTTGC TGACTAATTG AGATGCATGC TTTGCATACT TCTGCTGCT 300  
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255 ACGTCAATAA TGACGTATGT TCCCATAGTA ACGCCAATAG GGACTTTCCA TTGACGTCAA 540  
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**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/921,060B**

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**SEQUENCE MISSING ITEM REPORT**  
**PATENT APPLICATION US/08/921,060B**

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**SEQUENCE CORRECTION REPORT**  
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